Applicant: Ley et al. Attorney's Docket No.: 10280-074001 / YA-00-04

Serial No. : 09/884,767 Filed : June 19, 2001

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## Amendments to the Claims:

This listing of claims replaces all prior versions and listings of claims in the application:

## **Listing of Claims**:

- 1. (Currently amended) A non-naturally occurring enterokinase-cleavable fusion protein comprising a polypeptide comprising the formula:
  - (1)  $Z_1$ -Xaa<sub>1</sub>-Xaa<sub>2</sub>-Xaa<sub>3</sub>-Xaa<sub>4</sub>-Asp-Arg-Xaa<sub>5</sub>- $Z_2$  (SEQ ID NO:1), wherein
  - (a)  $Z_1$  is a ligand recognition sequence;
  - (b) Xaa<sub>1</sub>-Xaa<sub>2</sub>-Xaa<sub>3</sub>-Xaa<sub>4</sub>-Asp-Arg is an enterokinase recognition sequence, in which

Xaa<sub>1</sub> is Ala, Asp, Glu, Phe, Gly, Ile, Asn, Ser, or Val;

Xaa2 is Ala, Asp, Glu, His, Ile, Leu, Met, Gln or Ser;

Xaa<sub>3</sub> is Asp, Glu, Phe, His, Ile, Met, Asn, Pro, Val, or Trp, and

Xaa4- is Ala, Asp, Glu, or Thr; and

- (c)  $Xaa_5$ - $Z_2$  is a protein of interest, in which [[X5]]  $Xaa_5$  can be any amino acid and  $Z_2$  is a polypeptide of at least one amino acid.
  - 2. (Previously presented) The fusion protein of claim 1, wherein

Xaa<sub>1</sub> is Asp,

Xaa<sub>2</sub> is Ile,

Xaa<sub>3</sub> is Asn,

Xaa<sub>4</sub>-is Asp, and

Xaa<sub>5</sub>-is Met, Thr, Ser, Ala, Asp, Leu, Phe, Asn, Trp, Ile, Gln, Glu, His, Val, Gly or Tyr.

3. (Previously canceled)

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4. (Previously presented) The fusion protein of claim 1, wherein the ligand recognition sequence  $Z_1$  is a streptavidin binding domain.

5. (Original) The fusion protein of claim 4, wherein the streptavidin binding domain is selected from the sequences: His-Pro-Gln-Phe (SEQ ID NO:6), Cys-His-Pro-Gln-Phe-Cys (SEQ ID NO:5), Cys-His-Pro-Gln-Phe-Cys-Ser-Trp-Arg (SEQ ID NO:7), Trp-His-Pro-Gln-Phe-Ser-Ser (SEQ ID NO:210), Pro-Cys-His-Pro-Gln-Phe-Pro-Arg-Cys-Tyr (SEQ ID NO:211), and tandemly arranged combinations and repeats thereof.

## 6. -49. (Previously canceled)

- 50. (Previously presented) The fusion protein according to claim 1, wherein said ligand recognition sequence  $Z_1$  is selected from the group consisting of: streptavidin, avidin, an antibody, a peptide antigen recognized by the antibody, the Myc-tag, the Flag peptide, the KT3 epitope peptide, an  $\alpha$ -tubulin epitope peptide, a polyhistidine tag, a chitin binding domain, maltose binding protein (MBP), and the  $\alpha$  T7 gene 10-protein peptide tag.
- 51. (Previously presented) The fusion protein according to claim 1, wherein incubation of said polypeptide (SEQ ID NO:1) with enterokinase yields the protein of interest Xaa<sub>5</sub>-Z<sub>2</sub>.